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PC10960 Sequence Listing.ST25.txt  
SEQUENCE LISTING

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APR 04 2002

TECH CENTER 1600/2900

<110> Pfizer Ltd. (EP (GB) only)  
Pfizer Inc. (US, JP, EP except GB)  
Fidock, Mark David

<120> Novel Polypeptide

<130> PC10960AGPR

<150> GB 0030855.1

<151> 2000-12-18

<150> GB 0101222.8

<151> 2001-01-17

<160> 7

<170> PatentIn version 3.1

<210> 1

<211> 1082

<212> DNA

<213> Homo sapiens

<400> 1  
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ggtctctgcc agttctcaga gaagtacaag caagtctacc tctccctggc ctacagtatc 180  
atctttatcc tagggctgcc actaaatggc actgtcttgt ggcactcctg gggccaaacc 240  
aagcgctgga gctgtgccac cacctatctg gtgaacctga tgggtggccga cctgctttat 300  
gtgctattgc ccttctcat catcacctac tcactagatg acaggtggcc cttcggggag 360  
ctgctctgca agctggtgca cttcctgttc tatatcaacc ttacggcag catcctgctg 420  
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taccggaccc gcagggcatgc ctggctgggc accagcacca cctgggccct ggtggtcctc	540
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tatgacatga ccagccaaga gaattttgat cggcttttg cctacggcat agttctgaca	660
ttgtctggct ttctttccct ccttggtcat tttggtgtgc tattcactga tggtcaggag	720
cctgatcaag ccagaggaga acctcatgag gacaggcaac acagcccgag ccagggtccat	780
ccggaccatc ctactgggtgt gtggcctctt caccctctgt tttgtgccct tccatatcac	840
tcgctccttc tacctcacca tctgctttct gctttctcag gactgccagc tcttgatggc	900
agccagtgtg gcctacaaga tatggaggcc tctggtgagt gtgagcagct gcctcaaccc	960
agtcctgtac tttctttcaa ggggggcaaa aatagagtca ggctcctcca gaaactgagg	1020
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gg	1082

<210> 2

<211> 360

<212> PRT

<213> Homo sapiens

<400> 2

Met	Leu	Ser	Ile	Leu	Leu	Pro	Ser	Arg	Gly	Ser	Arg	Ser	Gly	Ser	Arg
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Arg	Gly	Ala	Leu	Leu	Leu	Glu	Gly	Ala	Ser	Arg	Asp	Met	Glu	Lys	Val
			20					25					30		
Asp	Met	Asn	Thr	Ser	Gln	Glu	Gln	Gly	Leu	Cys	Gln	Phe	Ser	Glu	Lys
		35					40					45			
Tyr	Lys	Gln	Val	Tyr	Leu	Ser	Leu	Ala	Tyr	Ser	Ile	Ile	Phe	Ile	Leu
	50					55					60				
Gly	Leu	Pro	Leu	Asn	Gly	Thr	Val	Leu	Trp	His	Ser	Trp	Gly	Gln	Thr
65					70					75					80
Lys	Arg	Trp	Ser	Cys	Ala	Thr	Thr	Tyr	Leu	Val	Asn	Leu	Met	Val	Ala
				85					90					95	
Asp	Leu	Leu	Tyr	Val	Leu	Leu	Pro	Phe	Leu	Ile	Ile	Thr	Tyr	Ser	Leu
			100					105					110		
Asp	Asp	Arg	Trp	Pro	Phe	Gly	Glu	Leu	Leu	Cys	Lys	Leu	Val	His	Phe
		115					120					125			

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Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile  
130 135 140

ser Val His Gln Phe Leu Gly Val Cys His Pro Leu Cys Ser Leu Pro  
145 150 155 160

Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala  
165 170 175

Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp  
180 185 190

Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn  
195 200 205

Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe  
210 215 220

Leu Ser Leu Leu Gly His Phe Gly Val Leu Phe Thr Asp Gly Gln Glu  
225 230 235 240

Pro Asp Gln Ala Arg Gly Glu Pro His Glu Asp Arg Gln His Ser Pro  
245 250 255

Ser Gln Val His Pro Asp His Pro Thr Gly Val Trp Pro Leu His Pro  
260 265 270

Leu Phe Cys Ala Leu Pro Tyr His Ser Leu Leu Leu Pro His His Leu  
275 280 285

Leu Ser Ala Phe Ser Gly Leu Pro Ala Leu Asp Gly Ser Gln Cys Gly  
290 295 300

Leu Gln Asp Met Glu Ala Ser Gly Glu Cys Glu Gln Leu Pro Gln Pro  
305 310 315 320

Ser Pro Val Leu Ser Phe Lys Gly Gly Lys Asn Arg Val Arg Leu Leu  
325 330 335

Gln Lys Leu Arg Gln Asn Lys Leu Gly Glu His Pro Ala Gly Arg Lys  
340 345 350

Arg Cys Pro Gly Leu Asn Arg Ser  
355 360

<210> 3

<211> 1020

<212> DNA

<213> Homo sapiens

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<400> 3
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ggtctctgcc agttctcaga gaagtacaag caagtctacc tctccctggc ctacagtatc      180
atctttatcc tagggctgcc actaaatggc actgtcttgt ggcactcctg gggccaaacc      240
aagcgctgga gctgtgccac cacctatctg gtgaacctga tgggtggccga cctgctttat      300
gtgctattgc ctttcctcat catcacctac tcactagatg acaggtggcc cttcggggag      360
ctgctctgca agctggtgca cttcctgttc tatatcaacc ttacggcag catcctgctg      420
ctgacctgca tctctgtgca ccagttccta ggtgtgtggc acccactgtg ttcgctgccc      480
taccggaccc gcaggcatgc ctggctgggc accagcacca cctgggccct ggtggtcctc      540
cagctgctgc ccacactggc cttctcccac acggactaca tcaatggcca gatgatctgg      600
tatgacatga ccagccaaga gaattttgat cggctttttg cctacggcat agttctgaca      660
ttgtctggct ttctttcccc ctccttggtc attttggtgt gctattcact gatggtcagg      720
agcctgatca agccagagga gaacctcatg aggacaggca acacagcccg agccagggtcc      780
atccggacca tcctactggt gtgtggcctc ttcaccctct gttttgtgcc cttccatatc      840
actcgctcct tctacctcac catctgcttt ctgctttctc aggactgcca gctcttgatg      900
gcaccagtg tggcctacaa gatatggagg cctctggtga gtgtgagcag ctgcctcaac      960
ccagtcctgt actttctttc aaggggggca aaaatagagt caggctcctc cagaaactga     1020

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<210> 4

<211> 339

<212> PRT

<213> Homo sapiens

<400> 4

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                20                25                30
Asp Met Asn Thr Ser Gln Glu Gln Gly Leu Cys Gln Phe Ser Glu Lys
          35                40                45
Tyr Lys Gln Val Tyr Leu Ser Leu Ala Tyr Ser Ile Ile Phe Ile Leu
    50                55                60

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Gly Leu Pro Leu Asn Gly Thr Val Leu Trp His Ser Trp Gly Gln Thr  
65 70 75 80

Lys Arg Trp Ser Cys Ala Thr Thr Tyr Leu Val Asn Leu Met Val Ala  
85 90 95

Asp Leu Leu Tyr Val Leu Leu Pro Phe Leu Ile Ile Thr Tyr Ser Leu  
100 105 110

Asp Asp Arg Trp Pro Phe Gly Glu Leu Leu Cys Lys Leu Val His Phe  
115 120 125

Leu Phe Tyr Ile Asn Leu Tyr Gly Ser Ile Leu Leu Leu Thr Cys Ile  
130 135 140

Ser Val His Gln Phe Leu Gly Val Trp His Pro Leu Cys Ser Leu Pro  
145 150 155 160

Tyr Arg Thr Arg Arg His Ala Trp Leu Gly Thr Ser Thr Thr Trp Ala  
165 170 175

Leu Val Val Leu Gln Leu Leu Pro Thr Leu Ala Phe Ser His Thr Asp  
180 185 190

Tyr Ile Asn Gly Gln Met Ile Trp Tyr Asp Met Thr Ser Gln Glu Asn  
195 200 205

Phe Asp Arg Leu Phe Ala Tyr Gly Ile Val Leu Thr Leu Ser Gly Phe  
210 215 220

Leu Ser Pro Ser Leu Val Ile Leu Val Cys Tyr Ser Leu Met Val Arg  
225 230 235 240

Ser Leu Ile Lys Pro Glu Glu Asn Leu Met Arg Thr Gly Asn Thr Ala  
245 250 255

Arg Ala Arg Ser Ile Arg Thr Ile Leu Leu Val Cys Gly Leu Phe Thr  
260 265 270

Leu Cys Phe Val Pro Phe His Ile Thr Arg Ser Phe Tyr Leu Thr Ile  
275 280 285

Cys Phe Leu Leu Ser Gln Asp Cys Gln Leu Leu Met Ala Pro Ser Val  
290 295 300

Ala Tyr Lys Ile Trp Arg Pro Leu Val Ser Val Ser Ser Cys Leu Asn  
305 310 315 320

Pro Val Leu Tyr Phe Leu Ser Arg Gly Ala Lys Ile Glu Ser Gly Ser  
325 330 335

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Ser Arg Asn

<210> 5

<211> 27

<212> DNA

<213> Homo sapiens

<400> 5

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27

<210> 6

<211> 24

<212> DNA

<213> Homo sapiens

<400> 6

tcaccagatc tggtcaaccc tggg

24

<210> 7

<211> 24

<212> DNA

<213> Homo sapiens

<400> 7

tcagtttctg gaggagcctg actc

24